Therefore, what I claim, is:

Claims:

- 1. A method for detecting resistant fungal cells in clinical material, comprising the steps of:
 - a) extraction of fungus-specific nucleic acids from clinical material; and
 - b) hybridization of the fungus-specific nucleic acids with hybridization probes which are directed against nucleic acid segments of azole derivative-resistant fungal cells.
- 2. The method as in Claim 1, wherein the hybridization probes are directed against a DNA segment from the $14-\alpha-$ lanosterol demethylase gene.
- 3. The method as in claim 2, wherein the hybridization probes are directed against a DNA segment from the $14-\alpha-$ lanosterol demethylase gene (ERG16 gene) of the species Candida albicans.
- 4. The method as in Claim 2, wherein between steps a) and b) a PCR reaction is performed in which segments of the $14-\alpha$ -lanosterol demethylase gene are amplified.

- The method as in Claim 3, wherein between steps a) and b) 5. a PCR reaction is performed in which segments of the $14-\alpha$ lanosterol demethylase gene are amplified.
- The method\as in Claim 4, wherein a primer for the PCR re-6. action is selected from the group consisting of SEQ ID-No.: 1, SEQ TO-No.: 2, SEQ ID-No.: 3 and SEQ ID-No.: 4.
- The method as ih Claim 5, wherein a primer for the PCR re-7. action is selected from the group consisting of SEQ ID-No.: 1, SEQ ID-No.: 2, SEQ IM-No.: 3 and SEQ ID-No.: 4.
- The method as in \Claim 1, wherein a hybridization probe 8. for step b) is selected from the group consisting of SEQ ID-No.: 5, SEQ ID-No.: 6, SEQ ID-No.: 7 and SEQ ID-No.:8.
- The method as in Claim 4, wherein a hybridization probe 9. for step b) is selected from the group consisting of SEQ ID-No.: 5, SEQ ID-No.: &, SEQ ID-No.: 7 and SEQ ID-No.:8.
- The method as in Claim 6, wherein a hybridization probe for step b) is selected from the group consisting of SEQ ID-No.: 5, SEQ ID-No\: 6, SEQ ID-No.: 7 and SEQ ID-No.:8.
 - The method as in Claim 1, wherein in step b) the hybridi-11. zation probes are labeled with digoxigenin and used in Southern hybridization.
- The method as in Claim 8, wherein after hybridization, at least one washing step is performed at a temperature which

is approximately 1°C less than the melting temperature (Tm) of the particular hybridization probe used.

- 13. The method as in Claim 9, wherein after hybridization, at least one washing step is performed at a temperature which is approximately 1°C less than the melting temperature (Tm) of the particular hybridization probe used.
- 14. The method as in Claim 10, wherein after hybridization, at least one washing step is performed at a temperature which is approximately 1°C less than the melting temperature (Tm) of the particular hybridization probe used.
- 15. The nucleotide sequence SEQ ID no. 1 from the enclosed Sequence Listing.
- 16. The nucleotide sequence SEQ ID no. 2 from the enclosed Sequence Listing.
- 17. The nucleotide sequence SEQ ID no. 3 from the enclosed Sequence Listing.
- 18. The nucleotide sequence SEQ ID no. 4 from the enclosed Sequence Listing.
- 19. The nucleotide sequence SEQ ID no. 5 from the enclosed Sequence Listing.
- 20. The nucleotide sequence SEQ ID no. 6 from the enclosed Sequence Listing.

- 21. The nucleotide sequence SEQ ID no. 7 from the enclosed Sequence Listing.
- 22. The nucleotide sequence SEQ ID no. 8 from the enclosed Sequence Listing.
- 23. Use of the nucleotide sequences SEQ ID no. 1 and SEQ ID no. 2 as primers, and nucleotide sequences SEQ ID no. 5 and/or SEQ ID no. 6 as hybridization probes, in a method as in Claim 4.
- 24. Use of the nucleotide sequences SEQ ID nos. 3 and SEQ ID no. 4 as primers, and nucleotide sequences SEQ ID no. 7 and/or SEQ ID no. 8 as hybridization probes, in a method as in Claim 4.
- 25. A kit for the analysis of fungal infections with azole derivative-resistant fungal strains, containing at least one nucleotide sequences selected from the group consisting of SEQ ID-No.: 1, SEQ ID-No.: 2, SEQ ID-No.: 3, SEQ ID-No.: 4, SEQ ID-No.: 5, SEQ ID-No.: 6, SEQ ID-No.: 7, SEQ ID-No.: 8.
- 26. A kit for performing the method as in Claim 1.

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